

FIGURE 1

TNIK 1 MASDSEAFSLDIDLSALRDPACIFELVGVNCTYGVYKGRHVKT COLAAIKVNDVTDEEEETKQETNMKKYSHHR
 NIK 1 MANDSEAFSLVDTLSSLRDPACIFELVGVNCTYGVYKGRHVKT-VTAALKVNDVTDEEEETKQETNMKKYSHHR

TNIK 81 HIATYYGAFIKKPEECDDQIWMVMEFFCAGSVTDIKNYKENTLKKEWIAPICREILRLSHLCHKVTHKIRGQNVL
 NIK 80 HIATYYGAFIKKPEECDDQIWMVMEFFCAGSVTDIKNYKENTLKKEWIAPICREILRLSHLCHKVTHKIRGQNVL

TNIK 161 LTENAEVKLVDFGVSAQLDRTVGRNRTFICTPYWMAPEVIACDENITATYDEKDLWSGITATFMAECAPPICHNHITR
 NIK 160 LTENAEVKLVDFGVSAQLDRTVGRNRTFICTPYWMAPEVIACDENITATYDEKDLWSGITATFMAECAPPICHNHITR

TNIK 241 ALFLIPRNPAPRLKSKKWSKKFSEFIESCIUVNHSQRPATEQIMKHITIRDOVNERQVHQLKDHIDRTKIKKGEKDETE
 NIK 240 ALFLIPRNPAPRLKSKKWSKKFSEFIESCIUVNHSQRPATEQIMKHITIRDOVNERQVHQLKDHIDRTKIKKGEKDETE

TNIK 321 YEYSGSEEEEREN--DSEPPSSINUGESTLRKDELRLLOENKERSEALKKQILOC--RENEEHKQOLLAERQKRIE
 NIK 320 YEYSGSEEEEREN--DSEPPSSINUGESTLRKDELRLLOENKERSEALKKQILOC--RENEEHKQOLLAERQKRIE

TNIK 397 EKEQKURLEEQQAEKELRSQEREOK-----NDEONSR--EERHNAHEEYTHQOLEEEOQOLE
 NIK 400 EKEQKURLEEQQAEKELRSQEREOK-----NDEONSR--EERHNAHEEYTHQOLEEEOQOLE

TNIK 460 ILQCOLLEONALLMYKFOLEKFAERLRLKQEROYLVSLQHQRQZQRPVKKIYHYNEGMSSEKFAWAKEVEE
 NIK 480 ILQCOLLEONALLMYKFOLEKFAERLRLKQEROYLVSLQHQRQZQRPVKKIYHYNEGMSSEKFAWAKEVEE

TNIK 540 RSLNROSSPAMTHKVANRISDPNTPERESEFISGVQPARTFPMINPVUQIHLVAVKSQGPALTASQSVHEQPTKGI
 NIK 532 RSLNROSSPAMTHKVANRISDPNTPERESEFISGVQPARTFPMINPVUQIHLVAVKSQGPALTASQSVHEQPTKGI

TNIK 620 SGFQALNVTSHRVEPRNSOPTSDNPELETRIEKFDRSSMLROEM--HUKVLCRTTSISEPAARKSPONCESALGPR
 NIK 584 S-----SSDSKSE-VPSE-OKAKSSSDSDEMPERVEVATTSRSIVILBANDSILQEGCOONS

TNIK 699 LSGPPIRANPDILRTBPIFESPTORTSSCS9SS9STPSSQSSQSGSFCSCAGSENNRVNANESKSGSPVLPHEPAK
 NIK 640 CAGRNSTSSIEPPLWERYEKLVRPSCSSSSCS9SS9SGSFCSCAGSENNRVNANESKSGSPVLPHEPAK

TNIK 779 VKPEESRDTTPSRPASVKKATOLULTALAKELRORLEETNOAKKVTIYSSSSEESSESEENICESETHNGAVAS
 NIK 716 VKPEESRDTTPSRPASVKKATOLULTALAKELRORLEETNOAKKVTIYSSSSEESSESEENICESETHNGAVAS

TNIK 859 DIPILPTICAPGNSQVNVGVCTIILETSHADSFCSISIEGTLMIETSGCKGSCSHDSNGFAGHINLPOLVQCSHG
 NIK 788 DIPILPTICAPGNSQVNVGVCTIILETSHADSFCSISIEGTLMIETSGCKGSCSHDSNGFAGHINLPOLVQCSHG

TNIK 939 PAGTPTGLGRVSTHSQEDSGCTEQCGSSKRASETPVPRVUTSUDDDEEDRESSAAALTSPLKCOLKNEAR
 NIK 846 -----SSSGFTEIDUPRLUISSS-----GTVVSEVVCSCDCLNFEALRODPTR

TNIK 1019 KISVVNVNPTNIREHSDTPEIRKYNKRNSEILCAALWGVNIVCTENHIMLDREGGKRVNINNNKNTLMVLEGLN
 NIK 892 KISVVNVNPTNIREHSDTPEIRKYNKRNSEILCAALWGVNIVCTENHIMLDREGGKRVNINNNKNTLMVLEGLN

TNIK 1099 VLVITISGKKOKLMVYYLSWLKNNILHNDPEVEKKQCMIVGULEGCTHYKVVKYERIKPIVJALKNAUETVYAWAPKIYHK
 NIK 972 VLVITISGKKOKLMVYYLSWLKNNILHNDPEVEKKQCMIVGULEGCTHYKVVKYERIKPIVJALKNAUETVYAWAPKIYHK

TNIK 1179 EMARKSEADLOIKPIILVLTVEEGORLKVIEGSHSTFIVDVUSCNSYDIYTHSHICNTIPIHATVILLVDEMEMLVY
 NIK 1052 EMARKSEADLOIKPIILVLTVEEGORLKVIEGSHSTFIVDVUSCNSYDIYTHSHICNTIPIHATVILLVDEMEMLVY

TNIK 1259 EDEGVYVNTYCRITKDVVLQWGEEMPTSVAYTHSNCTMGWGEKATEIRSVETGHLDCVEMHKRAQILKFLCANDKVFES
 NIK 1132 EDEGVYVNTYCRITKDVVLQWGEEMPTSVAYTHSNCTMGWGEKATEIRSVETGHLDCVEMHKRAQILKFLCANDKVFES

TNIK 1339 VRSGGSSQVEMTIAHNSMTN
 NIK 1212 VRSGGSSQVEMTIAHNSMTN

001200 001200 001200

FIGURE 2

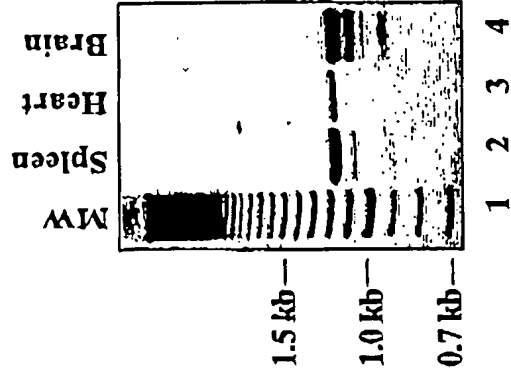


FIGURE 3

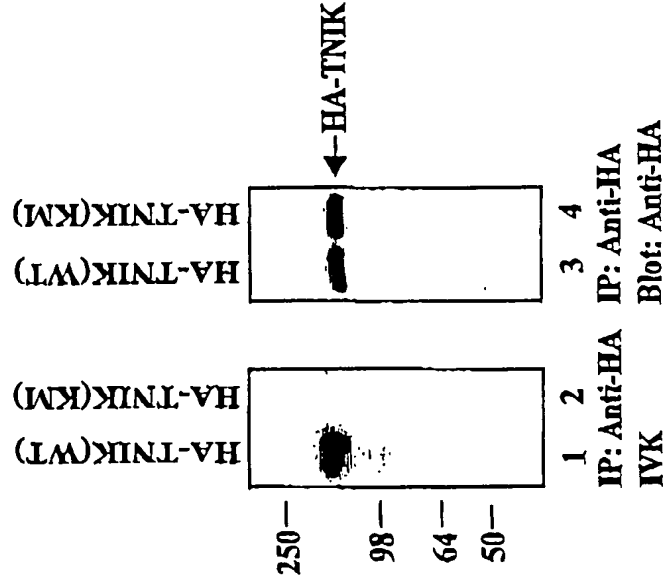
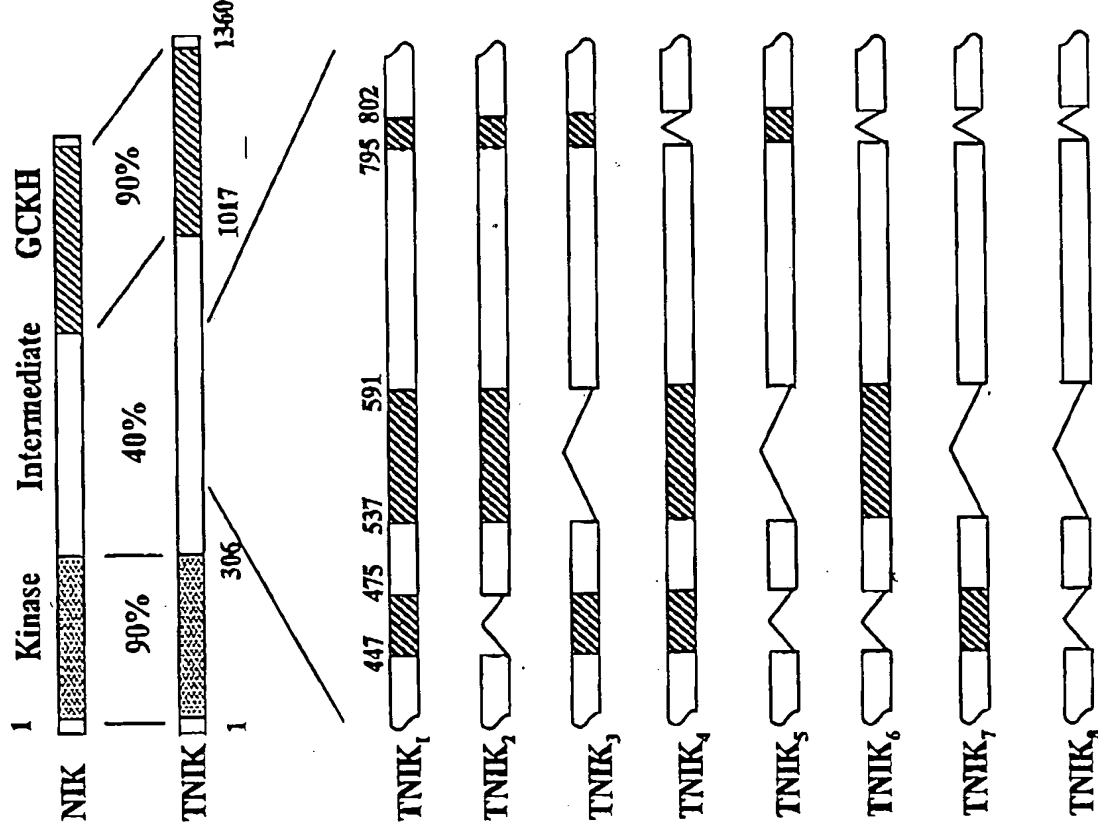


FIGURE 4

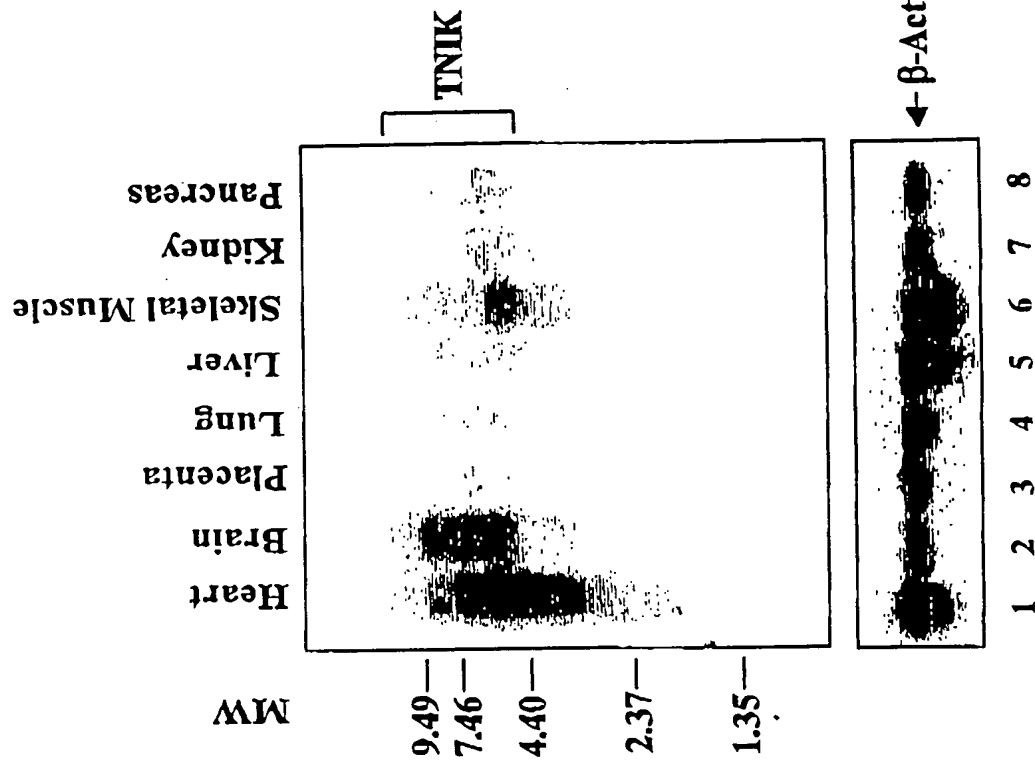


FIGURE 5A

FIGURE 5B

FIGURE 6

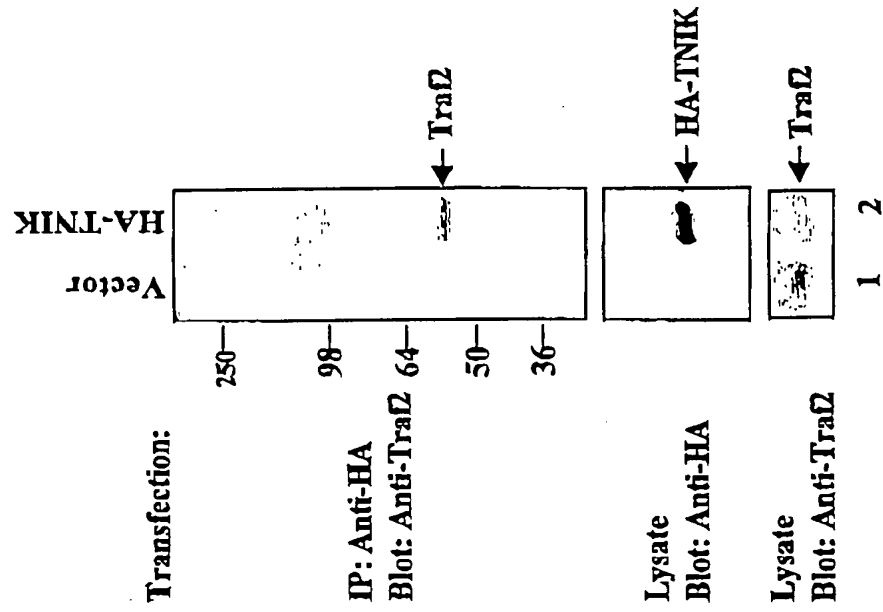


FIGURE 7

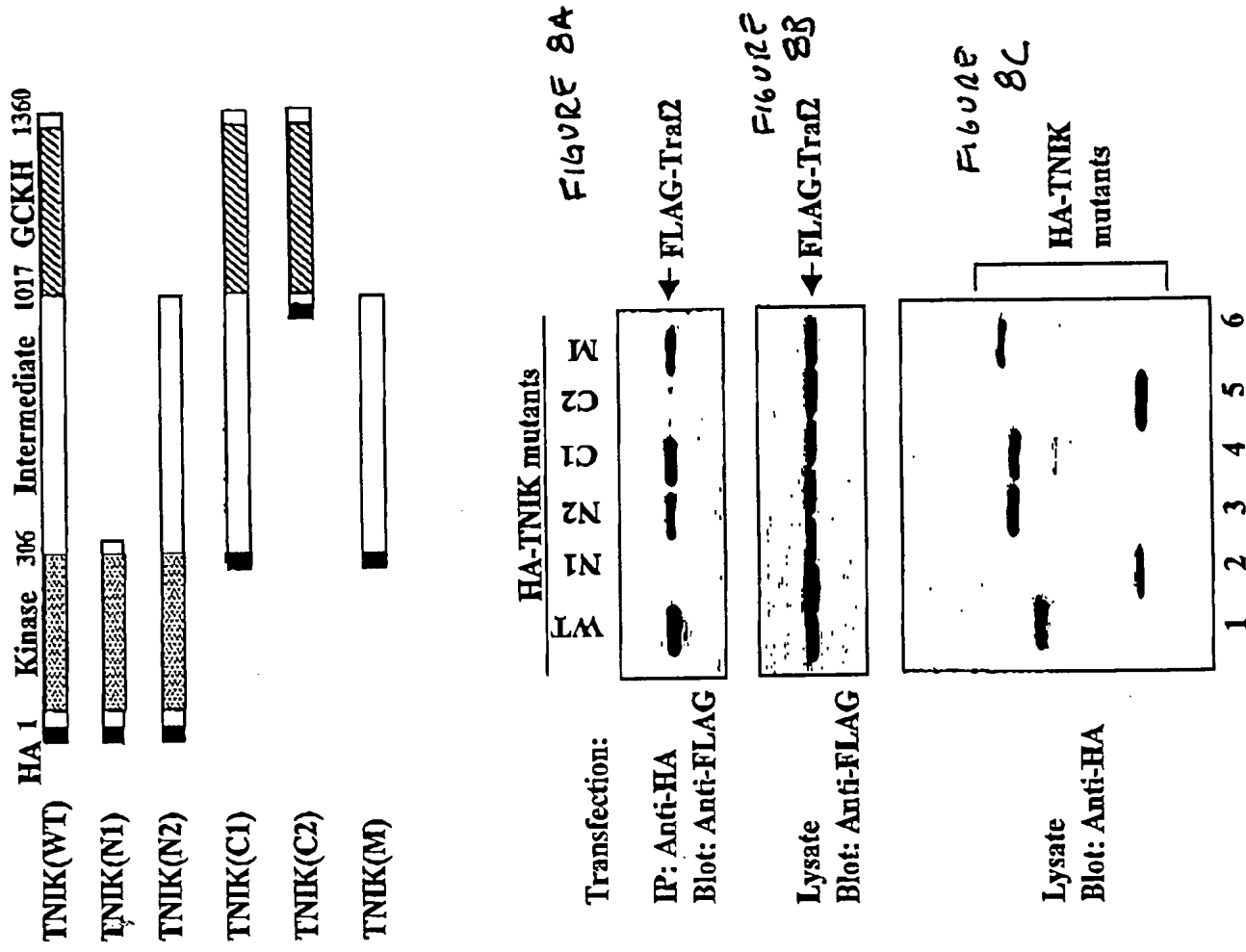


FIGURE 9

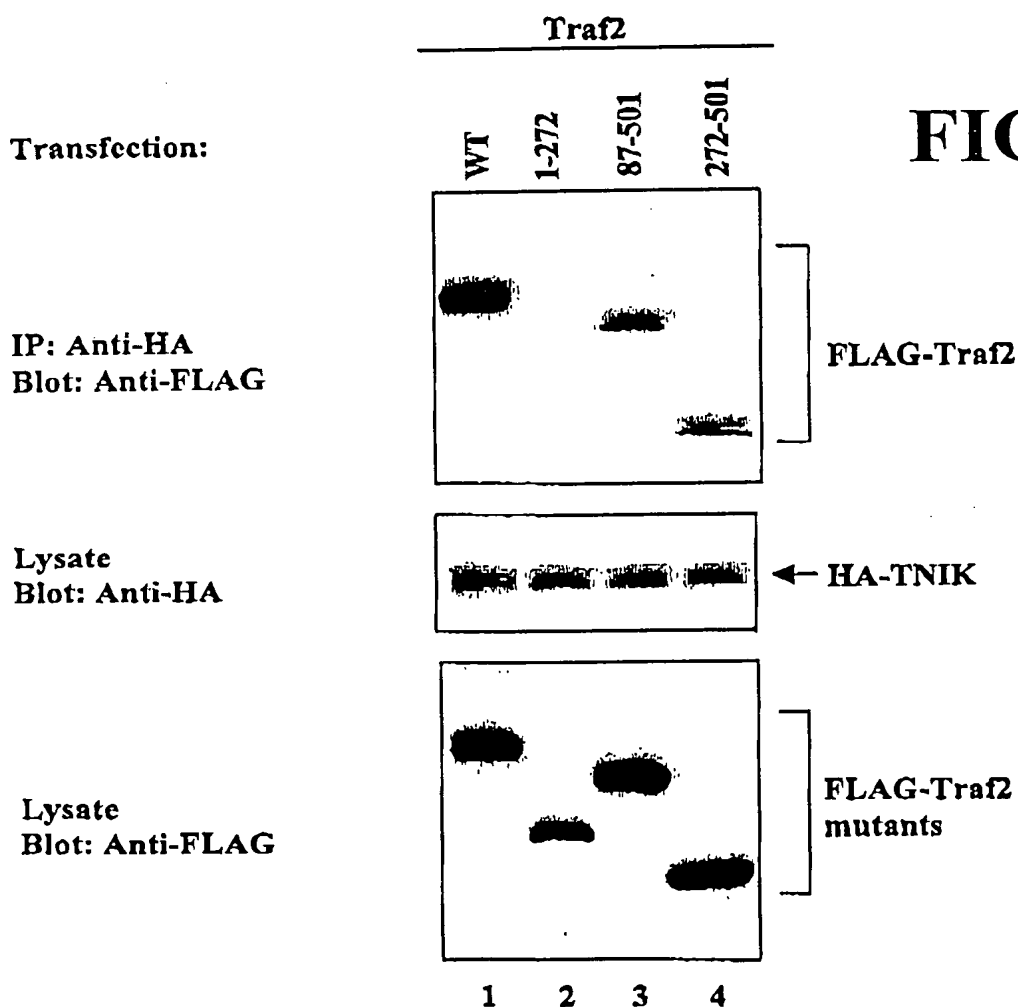
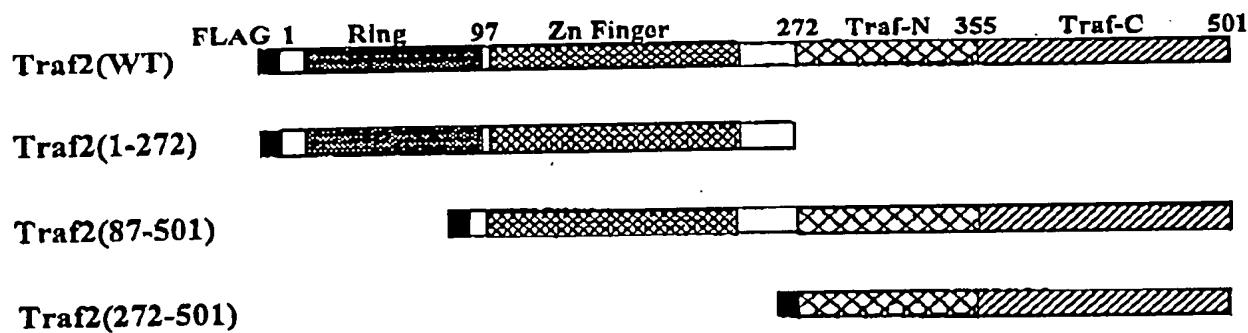


FIGURE 10

004280-95151560

FIGURE 11

B.

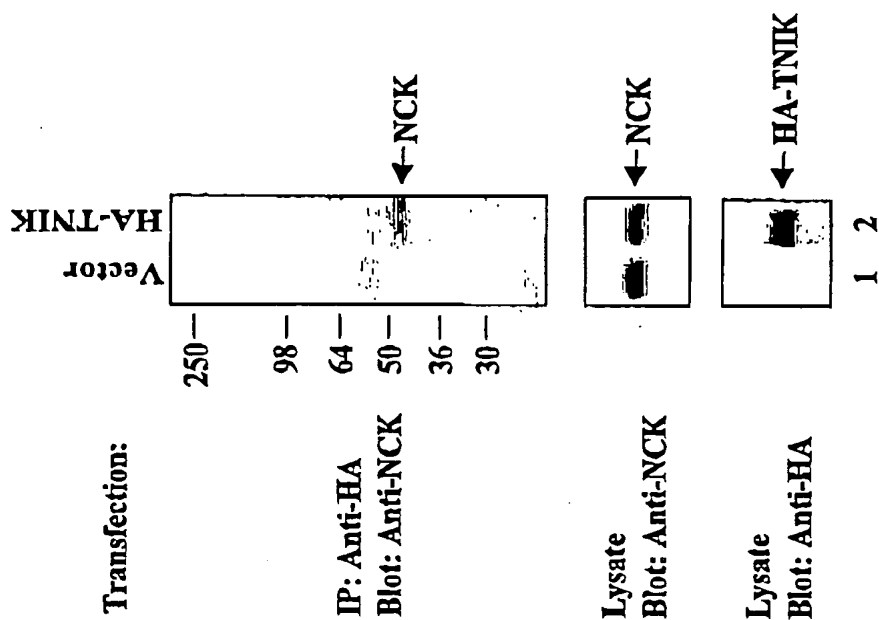


FIGURE 12

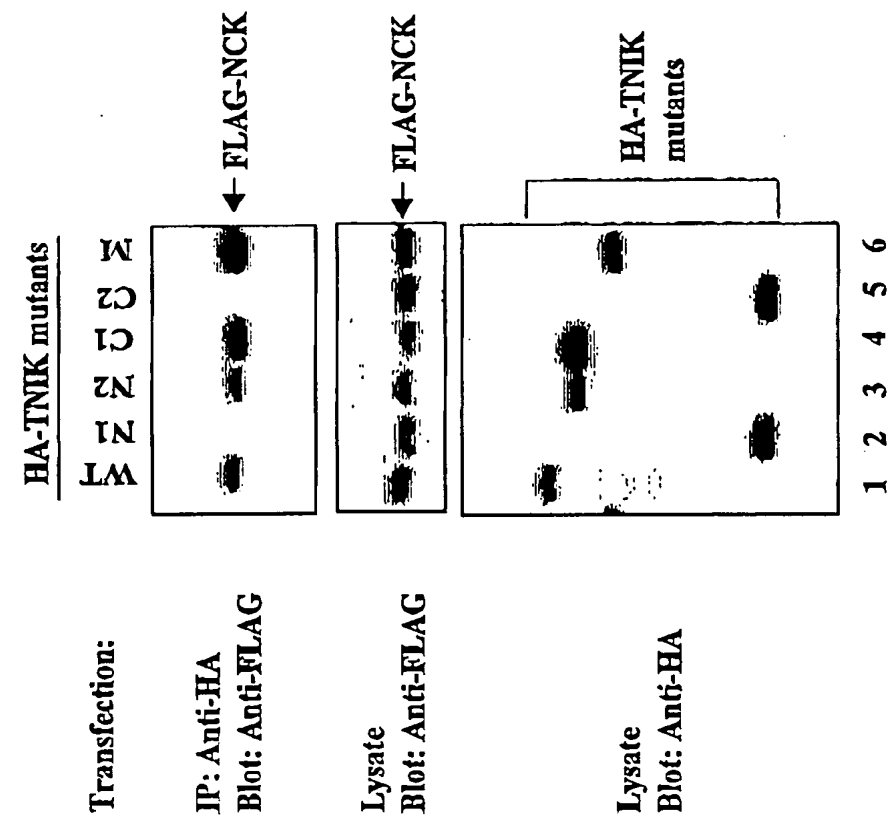
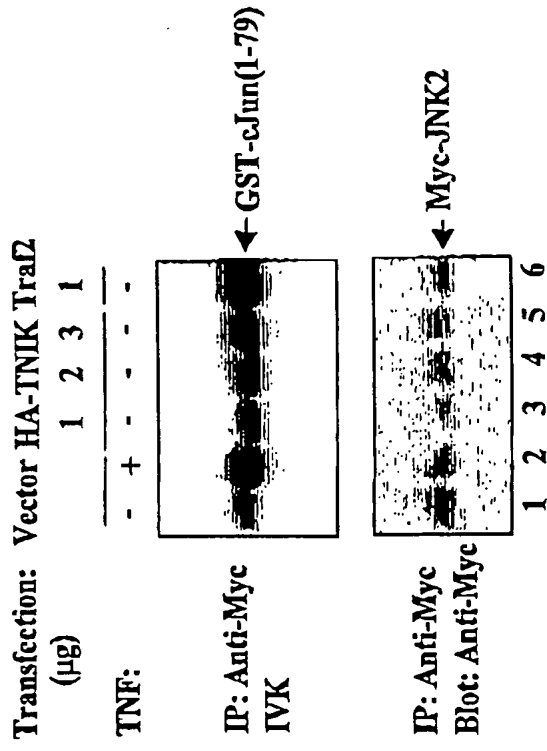


FIGURE 13



B.

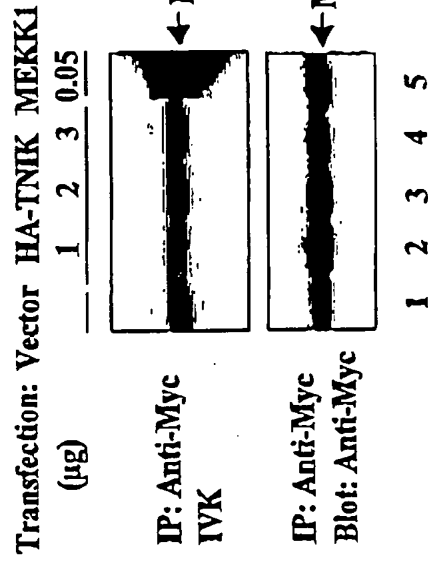


FIGURE 14

D.

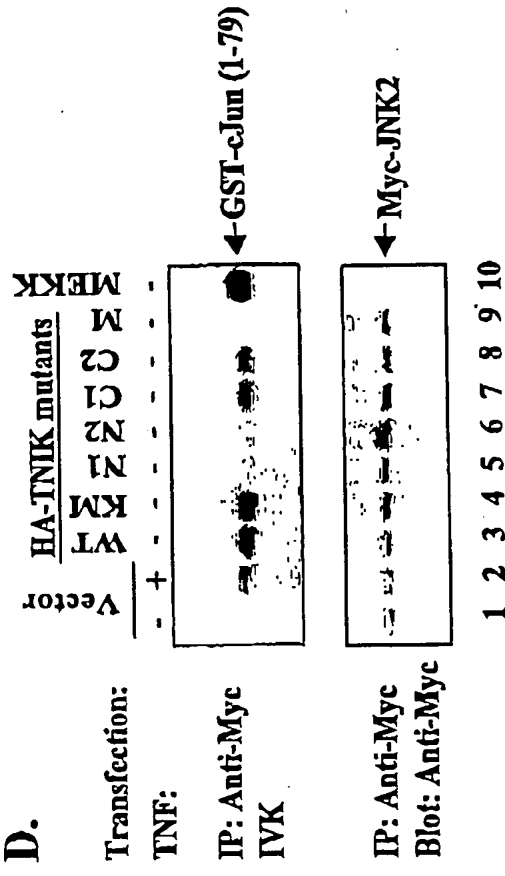
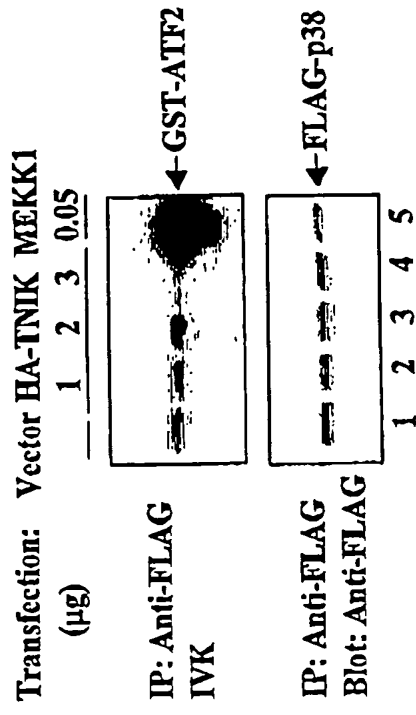


FIGURE 16

FIGURE 15



A.

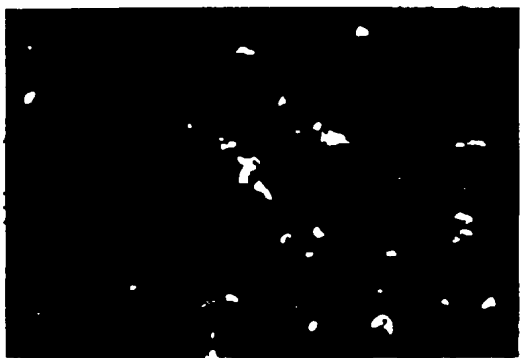
Vector



TNIK



TNIK(KM)



TNIK(N1)



TNIK(C1)



JNK2

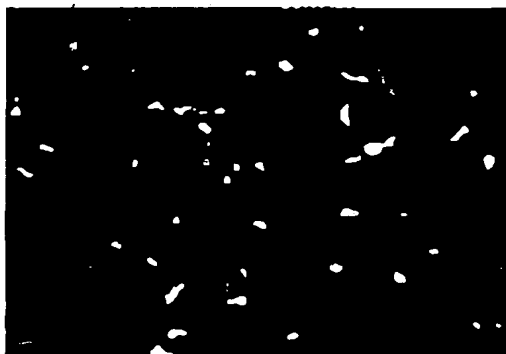
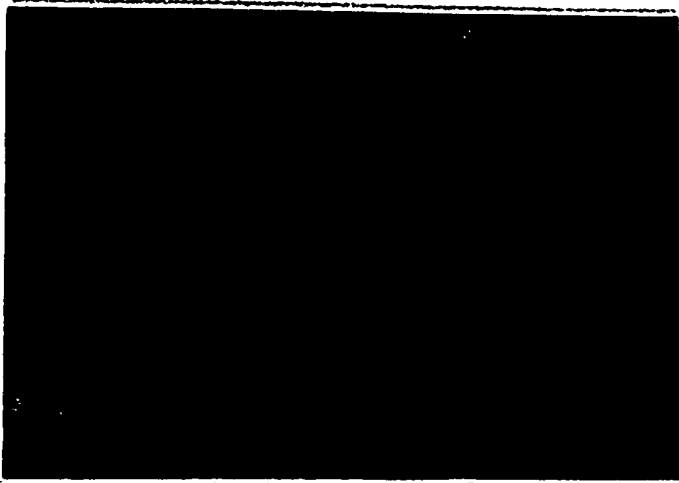


FIGURE 17

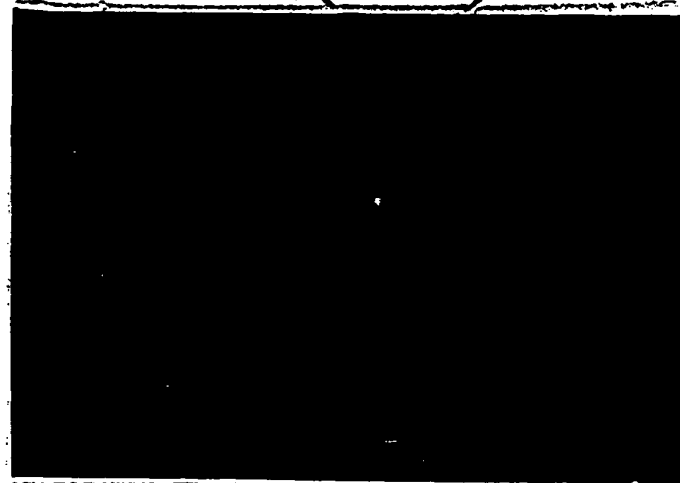
001230 93151530

FIGURE 18

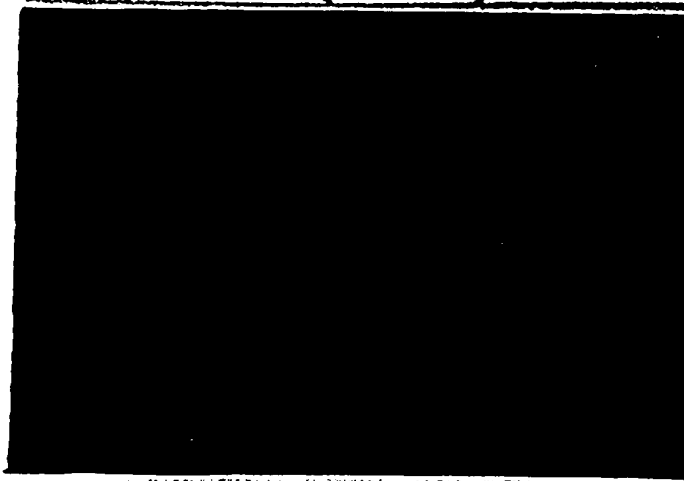
Vector



TNIK(WT)



TNIK(KM)



RIP

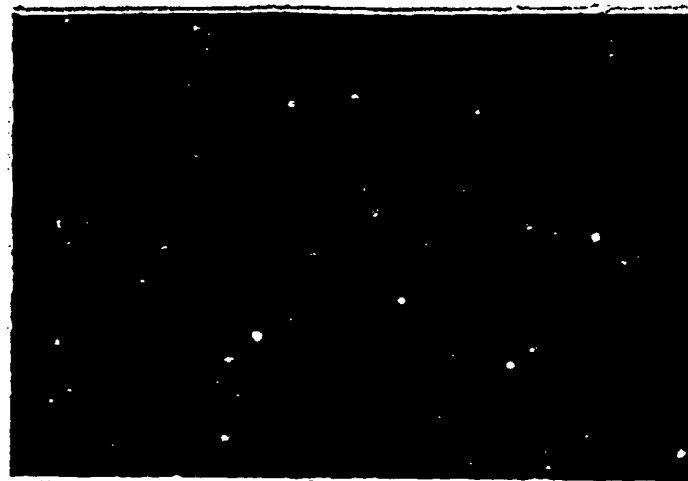


FIGURE 20

FIGURE 19

D.

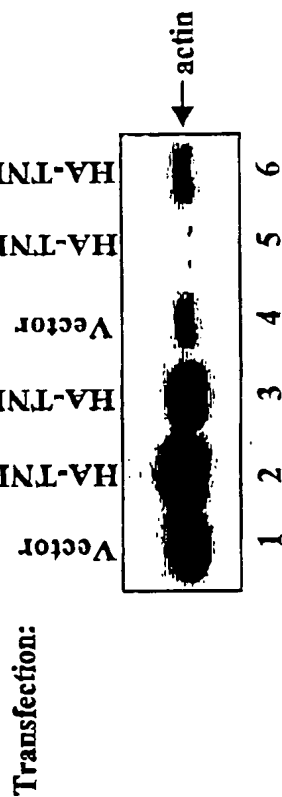
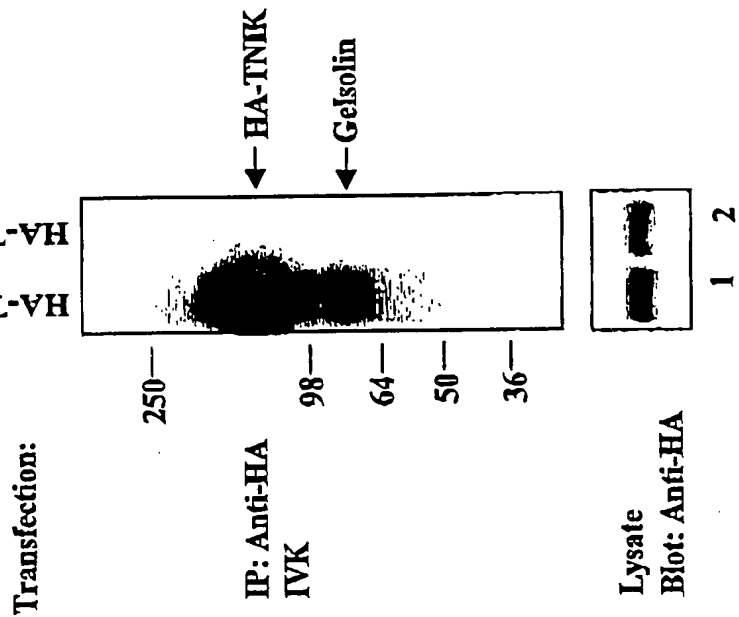


Figure 21

ATGGCGAGCGACTCCCCGGCTCGAAGCCTGGATGAAATAGATCTCTCGGCTCTGAGGGACCTGCAGGGATCTTT
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GCAGCCATCAAGGTTATGGATGTACAGGGGATGAAGAGGAAGAAATCAAAACAAGAAATTAACATGTTGAAGAAA
TATTCTCATCACCGGAATATTGCTACATACTATGGTGCTTTTATCAAAAAGAACCACCAGGCATGGATGACCAA
CTTTGGTTGGTGATGGAGTTTTGTGGTGCTGGCTCTGTACCGACCTGATCAAGAACACAAAAGGTAAACAGTTG
AAAGAGGAGTGGATTGCATACATCTGCAGGGAAATCTTACGGGGGCTGAGTCACCTGCACCAGCATAAAGTGATT
CATCGAGATATTAAAGGGCAAAATGTCTTGCTGACTGAAAATGCAGAAGTTAACTAGTGGACTTTGGAGTCAGT
GCTCAGCTTGATCGAACAGTGGGCAGGAGGAATACTTTCATTGGAACCTCCTACTGGATGGCACCAGAAGTTATT
GCCTGTGATGAAAACCCAGATGCCACATATGATTTCAAGAGTGACTTGTGGTCTTTGGGTATCACCGCCATTGAA
ATGGCAGAAGGTGCTCCCCCTCTCTGTGACATGCACCCCATGAGAGCTCTCTTCCTCATCCCCCGAATCCAGCG
CCTCGGCTGAAGTCTAAGAAGTGGTCAAAAAAATTCAGTCATTTATTGAGAGCTGCTTGGTAAAGAATCACAGC
CAGCGACCAGCAACAGAACAAATTGATGAAGCATCCATTTATACGAGACCAACCTAATGAGCGACAGGTCCGCATT
CAACTCAAGGACCATATTGATAGAACAAGAAGAAAGCGAGGAGAAAAAGATGAGACAGAGTATGAGTACAGTGGA
AGTGAGGAAGAAGAGGAGGAGAATGACTCAGGAGAGCCAGCTCCATCCTGAATCTGCCAGGGGAGTCGACGCTG
CGGAGGGACTTTCTGAGGCTGCAGCTGGCCAACAAGGAGCGTTCTGAGGCCCTACGGAGGCAGCAGCTGGAGCAG
CAGCAGCGGGAGAATGAGGAGCACAAAGCGGCAGCTGCTGGCCGAGCGTCAGAAGCGCATCGAGGAGCAGAAAGAG
CAGAGGCGGCGGCTGGAGGAGCAACAAGGCGAGAGAAGGAGCTGCGGAAGCAGCAGGAGAGGGAGCAGCGCCGG
CACTATGAGGAGCAGATGCGCCGGGAGGAGGAGAGGAGGCGTGCGGAGCATGAACAGGAATACATCAGGCGACAG
TTAGAGGAGGAGCAGAGACAGTTAGAGATCTTGACGACGAGCTACTGCATGAACAAGCTCTACTTCTGGAATAT
AAGCGCAAACAATTGGAAGAACAGAGACAAGCAGAAAGACTGCAGAGGCAGCTAAAGCAAGAAAGAGACTACTTA
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AGTCCTAGTGAGAAGCCAGCATGGGCCAAGGAGGTAGAAGAACGGTCAAGGCTCAACCGGCAAGTTCCCTGCC
ATGCCTCACAAGGTTGCCAACAGGATATCTGACCCCAACCTGCCCCCAAGGTGCGAGTCCTTCAGCATTAGTGGA
GTTACAGCTGCTCGAACACCCCCCATGCTCAGACCAGTCGATCCCCAGATCCCACATCTGGTAGCTGTAAAATCC
CAGGGACCTGCCTTGACCGCCTCCAGTCAGTGACGAGCAGCCCACAAAGGGCCTCTCTGGGTTTCAGGAGGCT
CTGAACGTGACCTCCACCGCGTGGAGATGCCACGCCAGAACTCAGATCCCACCTCGGAAAATCCTCCTCTCCCC
ACTCGCATTGAAAAGTTTGACCGAAGCTCTTGTTACGACAGGAAGAAGACATTCCACCAAAGGTGCCTCAAAGA
ACAACCTCTATATCCCCAGCATTAGCCAGAAAGAATTCTCCTGGGAATGGTAGTGCTCTGGGACCCAGCATAGGA
TCTCAACCCATCAGAGCAAGCAACCCTGATCTCCGGAGAACTCAGCCCATCTTGGAGAGCCCCCTGCAGAGGACC
AGCAGTGCGAGTTCTCTCAGCTCCAGCACCCCTAGCTCCAGCCAGCTCCCAAGGAGGCTCCAGCCTGGATCA
CAAGCAGGATCCAGTACAGCACAGAGTTTCGAGGCCAACAGTAAGTCAGAAGGATCACCTGTGCTTCCCCATGAG
CCTGCCAAGGTGAAACCAGAAAGAAATCCAGGGACATTACCCGGCCCCAGTCGACCAGCTAGCTACAAAAAGCTATA
GATGAGGATCTGACGGCATTAGCCAAAGAACTAAGAGAATCCGGATTGAAGAAAACAAACCGCCCAATGAAGAAG
GTGACTGATTACTCCTCCTCCAGTGAGGAGTCAGAAAGTAGCGAGGAAGAGGAGGAAGATGGAGAGAGCGAGACC
CATGATGGGACAGTGGCTGTGACGACATACCCAGACTGATACCAACAGGAGCTCCAGGCAGCAACGAGCAGTAC
AATGTGGGAATGGTGGGGACGCATGGGCTGGAGACCTCTCATGCGGACAGTTTCAGCGGCAGTATTTCAAGAGAA
GGAACCTTGATGATTAGAGAGACGTCTGGAGAGAAGAAGCGATCTGGCCACAGTGACAGCAATGGCTTTGTGGC
CACATCAACCTCCCTGACCTGGTGCAGCAGAGCCATTCTCCAGCTGGAACCCGACTGAGGGACTGGGGCGCGTC
TCAACCCATTCCAGGAGATGGACTCTGGGACTGAATATGGCATGGGGAGCAGCACCAAGCCTCCTTCACCCCC
TTTGTGGACCCAGAGTATACCAGACGTCTCCCACTGATGAAGATGAAGAGGATGAGGAATCATCAGCCGCAGCT
CTGTTTACTAGCGAACTTCTTAGGCAAGAACAGGCCAAACTCAATGAAGCAAGAAAGATTTCCGGTGGTAAATGTA
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CTTTGTGCAGCTCTGTGGGGTGTAACCTTCTGGTGGGGACTGAAAATGGCCCTGATGCTTTTGGACCGAAGTGGG
CAAGGCAAAGTCTATAATCTGATCAACCGGAGGCGATTTTCAGCAGATGGATGTGCTAGAGGGACTGAATGTCTT
GTGACAATTTTCAGGAAAGAAGAATAAGCTACGAGTTTACTATCTTTCATGGTTAAGAAACAGAATACTACATAAT
GACCCAGAAGTAGAAAAGAAACAAGGCTGGATCACTGTTGGGGACTTGGAAGGCTGTATACATTATAAAGTTGTT
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TATCATAAATTCATGGCATTAAAGTCTTTTGACAGATCTCCAGCACAAAGCCTCTGCTAGTTGATCTCAGCGTAGAA
GAAGGTCAAAGATTAAGGTTATTTTTGGTTACACACTGCTTCCATGTAATTGATGTTGATTTCAGGAACTCT
TATGATATCTACATACCATCTCATATTCAGGGCAATATCATCTCCATGCTATTGTCTATCTTGCTTAAACAGAT
GGAATGGAATGCTTGTGCTATGAGGATGAGGGGGTGATGTAAACACCTATGGCCGGATAACTAAGGATGTG
GTGCTCCAATGGGGAGAAATGCCACGTCTGTGGCCTACATTCATTCCAATCAGATAATGGGCTGGGGCGAGAAA
GCTATTGAGATCCGGTCAGTGGAACAGGACATTTGGATGGAGTATTTATGCATAAGCGAGCTCAAAGGTTAAAG
TTTCTATGTGAAAGAAATGATAAGGTATTTTTTGCATCCGTGCGATCTGGAGGAAGTAGCCAAGTGTTTTTCATG
ACCCTCAACAGAAATTCATGATGAACTGGTAA

Variable	Mean	SD	Min	Max
Age	38.5	12.5	18	65
Gender	0.5	0.5	0	1
Marital status	0.7	0.5	0	1
Education	12.5	2.5	9	16
Income	1500	500	500	3000
Health status	0.8	0.4	0	1
Smoking status	0.3	0.5	0	1
Alcohol consumption	0.2	0.4	0	1
Exercise frequency	0.5	0.5	0	1
Stress level	0.6	0.5	0	1
Depression score	0.4	0.5	0	1
Life satisfaction	0.7	0.5	0	1
Work satisfaction	0.6	0.5	0	1
Family satisfaction	0.7	0.5	0	1
Community satisfaction	0.6	0.5	0	1
Overall well-being	0.6	0.5	0	1

Variable	Mean	SD	Min	Max
Age	38.5	12.5	18	65
Gender	0.5	0.5	0	1
Marital status	0.7	0.5	0	1
Education	12.5	2.5	9	16
Income	1500	500	500	3000
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Alcohol consumption	0.2	0.4	0	1
Exercise frequency	0.5	0.5	0	1
Stress level	0.6	0.5	0	1
Depression score	0.4	0.5	0	1
Life satisfaction	0.7	0.5	0	1
Quality of life	0.8	0.4	0	1
Healthcare utilization	0.6	0.5	0	1
Health insurance status	0.9	0.3	0	1
Healthcare access	0.7	0.5	0	1
Healthcare cost	1000	300	500	2000
Healthcare quality	0.8	0.4	0	1
Healthcare satisfaction	0.7	0.5	0	1
Healthcare utilization frequency	0.5	0.5	0	1
Healthcare utilization cost	500	200	250	1000
Healthcare utilization quality	0.8	0.4	0	1
Healthcare utilization satisfaction	0.7	0.5	0	1
Healthcare utilization frequency cost	250	100	125	500
Healthcare utilization frequency quality	0.8	0.4	0	1
Healthcare utilization frequency satisfaction	0.7	0.5	0	1
Healthcare utilization cost quality	0.8	0.4	0	1
Healthcare utilization cost satisfaction	0.7	0.5	0	1
Healthcare utilization quality satisfaction	0.8	0.4	0	1
Healthcare utilization frequency cost quality	0.8	0.4	0	1
Healthcare utilization frequency cost satisfaction	0.7	0.5	0	1
Healthcare utilization frequency quality satisfaction	0.8	0.4	0	1
Healthcare utilization cost quality satisfaction	0.8	0.4	0	1
Healthcare utilization cost frequency satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency satisfaction	0.8	0.4	0	1
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Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization frequency quality cost satisfaction	0.8	0.4	0	1
Healthcare utilization frequency quality cost satisfaction	0.7	0.5	0	1
Healthcare utilization cost frequency quality satisfaction	0.8	0.4	0	1
Healthcare utilization cost frequency quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
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Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
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Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
Healthcare utilization frequency cost quality satisfaction	0.8	0.4	0	1
Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
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Healthcare utilization frequency cost quality satisfaction	0.8	0.4	0	1
Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
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Healthcare utilization frequency cost quality satisfaction	0.8	0.4	0	1
Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
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Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
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Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
Healthcare utilization frequency cost quality satisfaction	0.8	0.4	0	1
Healthcare utilization frequency cost quality satisfaction	0.7	0.5	0	1
Healthcare utilization quality frequency cost satisfaction	0.8	0.4	0	1
Healthcare utilization quality frequency cost satisfaction	0.7	0.5	0	1
Healthcare utilization frequency cost quality satisfaction	0.			

Figure 23

ATGGCGAGCGACTCCCCGGCTCGAAGCCTGGATGAAATAGATCTCTCGGCTCTGAGGGACCCTGCAGGGATCTTT
GAATTGGTGGAACTTGTGGAAATGGAACATACGGGCAAGTTTATAAGGGTCGTATGTCAAAACGGGCCAGCTT
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TATTCTCATCACCGGAATATTGCTACATACTATGGTGCTTTTATCAAAAAGAACCCACCAGGCATGGATGACCAA
CTTTGGTTGGTGTATGGAGTTTTTGTGGTGCTGGCTCTGTACCGACCTGATCAAGAACACAAAAGGTAACACGTTG
AAAGAGGAGTGGATTGCATACATCTGCAGGGAAATCTTACGGGGGCTGAGTCACCTGCACCAGCATAAAGTGATT
CATCGAGATATTAAAGGGCAAATGTCTTGCTGACTGAAAATGCAGAAGTTAAACTAGTGGACTTTGGAGTCAGT
GCTCAGCTTGATCGAACAGTGGGCAGGAGGAATACTTTTCAATGGAACCTCCTACTGGATGGCACCAGAAGTTATT
GCCTGTGATGAAAACCCAGATGCCACATATGATTTCAAGAGTGACTTGTGGTCTTTGGGTATCACCGCCATTGAA
ATGGCAGAAGGTGCTCCCCCTCTCTGTGACATGCACCCCATGAGAGCTCTCTTCTCATCCCCCGAATCCAGCG
CCTCGGCTGAAGTCTAAGAAGTGGTCAAAAAAATTCAGTCATTTATTGAGAGCTGCTTGGTAAAGAATCACAGC
CAGCGACCAGCAACAGAACAATTGATGAAGCATCCATTTATACGAGACCAACCTAATGAGCGACAGGTCCGCATT
CAACTCAAGGACCATATTGATAGAACAAGAAGAAGCGAGGAGAAAAAGATGAGACAGAGTATGAGTACAGTGGA
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CAGCAGCGGGAGAATGAGGAGCACAAAGCGGCAGCTGCTGGCCGAGCTCAGAAGCGCATCGAGAGCGAGAAAGAG
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Figure 24

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Figure 25

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GAACGCACCAGAGTTCGAGCCAACAGTAAGTCAGAAGGATCACCTGTGCTTCCCATGAGCCTGCCAAGGTGAAA
CCAGAAGAATCCAGGGACATTACCCGGCCCCAGTCGACCAGCTGATCTGACGGCATTAGCCAAAGAAGTAAGAGAA
CTCCGGATTGAAGAAACAACCCGCCAATGAAGAAGGTGACTGATTACTCTCTCCCTCCAGTGAGGAGTCAGAAAGT
AGCGAGGAAGAGGAGGAGAATGGAGAGGAGCAGCAAGCCATGATGGGACAGTGGCTGTGACGCACATACCCAGACTG
ATACCAACAGGAGCTCCAGGCAGCAACGAGCAGTACAATGTGGGAATGGTGGGGACGCATGGGCTGGAGACCTCT
CATGCGGACAGTTTCAGCGGCAGTATTTCAAGAGAAGGAACCTTGATGATTAGAGAGACGTCTGGAGAGAAGAAG
CGATCTGGCCACAGTGACAGCAATGGCTTTGCTGGCCACATCAACCTCCCTGACCTGGTGCAGCAGAGCCATTCT
CCAGCTGGAACCCCGACTGAGGGACTGGGGCGCGTCTCAACCCATTCCAGGAGATGGACTCTGGGACTGAATAT
GGCATGGGGAGCAGCACCAAAGCCTCCTTCACCCCTTTGTGGACCCAGAGTATACCAGACGTCTCCCACTGAT
GAAGATGAAGAGGATGAGGAATCATCAGCCGAGCTCTGTTTACTAGCGAACTTCTTAGGCAAGAACAGGCCAAA
CTCAATGAAGCAAGAAAGATTTGGTGGTAAATGTAAACCAACCAACATTTCGGCCTCATAGCGACACACCAGAA
ATCAGAAAATACAAGAAACGATTCAACTCAGAAATACTTTGTGCAGCTCTGTGGGGTGTAACCTTCTGGTGGGG
ACTGAAAATGGCCTGATGCTTTTGGACCGAAGTGGGCAAGGCAAAGTCTATAATCTGATCAACCGGAGGCGATT
CAGCAGATGGATGTGCTAGAGGGACTGAATGTCCTTGTGACAATTTTCAGGAAAGAAGAATAAGCTACGAGTTTAC
TATCTTTTCATGGTTAAGAAACAGAATACTACATAATGACCCAGAAGTAGAAAAAGAAACAAGGCTGGATCACTGTT
GGGGACTTGGAAGGCTGTATACATTATAAAGTTGTTAAATATGAAAGGATCAAATTTTTGGTGATTGCCTTAAAG
AATGCTGTGGAAATATATGCTTGGGCTCCTAAACCGTATCATAAATTCATGGCATTAAAGTCTTTTGCAGATCTC
CAGCACAAGCCTCTGCTAGTTGATCTCAGCGTAGAAGAAGGTCAAAGATTAAAGGTTATTTTTGGTTCACACACT
GGTTTCCATGTAATTGATGTTGATTTCAGGAACTCTTATGATATCTACATACCATCTCATATTCAGGGCAATATC
ACTCCTCATGCTATTGTCTGCTTGCCATAAACAGATGGAATGGAAATGCTTGTGCTATGAGGATGAGGGGGTG
TATGTAACACCTATGGCCCGGATAACTAAGGATGTGGTGCTCCAATGGGGAGAAATGCCACGCTGTGGCCATAC
ATTCAATTCCAATCAGATAATGGGCTGGGGCGAGAAAGCTATTGAGATCCGGTCAGTGGAACACAGACATTGGAT
GGAGTATTTATGCATAAGCGAGCTCAAAGGTTAAAGTTTCTATGTGAAAGAAATGATAAGGTATTTTTTGCATCC
GTGCGATCTGGAGGAAGTAGCCAAGTGTTTTTTCATGACCTTCAACAGAAATTCATGATGAACCTGGTAA

Figure 28

ATGGCGAGCGACTCCCCGGCTCGAAGCCTGGATGAAATAGATCTCTCGGCTCTGAGGGACCTGCAGGGATCTTT
GAATTGGTGGAACTTGTGGAAATGGAACATACGGGCAAGTTTATAAGGGTCGTATGTCAAAACGGGCCAGCTT
GCAGCCATCAAGGTTATGGATGTCACAGGGGATGAAGAGGAAGAAATCAAACAAGAAATTAACATGTTGAAGAAA
TATTCTCATCACCGGAATATTGCTACATACTATGGTGCCTTTTATCAAAAAGAACCACCAGGCATGGATGACCAA
CTTTGGTTGGTGTATGGAGTTTTTGTGGTGCTGGCTCTGTCACCGACCTGATCAAGAACACAAAAGGTAACACGTTG
AAAGAGGAGTGGATTGCATACATCTGCAGGGGAAATCTTACGGGGGCTGAGTCACCTGCACCAGCATAAAGTGATT
CATCGAGATATTAAAGGGCAAATGTCTTGCTGACTGAAAATGCAGAAAGTTAAACTAGTGGAAGTTTGGAGTCAGT
GCTCAGCTTGATCGAACAGTGGGCAGGAGGAATACTTTCATTGGAAGTCCCTACTGGATGGCACCAGAAGTTATT
GCCTGTGATGAAAACCCAGATGCCACATATGATTTCAAGAGTGACTTGTGGTCTTTGGGTATCACCGCCATTGAA
ATGGCAGAAGGTGCTCCCCCTCTCTGTGACATGCACCCCATGAGAGCTCTCTTCTCATCCCCCGGAATCCAGCG
CCTCGGCTGAAGTCTAAGAAGTGGTCAAAAAAATTCAGTCATTTATTGAGAGCTGCTTGGTAAAGAATCACAGC
CAGCGACCAGCAACAGAACAAATTGATGAAGCATCCATTTATACGAGACCAACCTAATGAGCGACAGGTCCGCTT
CAACTCAAGGACCATATTGATAGAACAAGAAGAAGCGAGGAGAAAAAGATGAGACAGAGTATGAGTACAGTGGA
AGTGAGGAAGAAGAGGAGGAGAATGACTCAGGAGAGCCAGCTCCATCTGAATCTGCCAGGGGAGTCGACGCTG
CGGAGGGACTTTCTGAGGCTGCAGCTGGCCAAACAAGGAGCGTTCGAGGCCCTACGGAGGCAGAGCTGGAGCAG
CAGCAGCGGGAGAATTGAGGAGCACAAAGCGGCAGCTGCTGGCCGAGCGTCAGAAGCGCATCGAGGAGCAGAAAGAG
CAGAGGCGGCGGCTGGAGGAGCAACAAAGGCGAGAGAAGGAGCTGCGGAAGCAGCAGGAGAGGGAGCAGCGCCGG
CACTATGAGGAGCAGATGCGCCGGGAGGAGGAGGAGGAGGCGTGCAGGAGCATGAACAGGAATATAAGCGCAAACAA
TTGGAAGAACAGAGACAAGCAGAAAGACTGCAGAGGCAGCTAAAGCAAGAAAGAGACTACTTAGTTTCCCTTCAG
CATCAGCGGCAGGAGCAGAGGCCTGTGGAGAAGAAGCCACTGTACCATTACAAAGAAGGAATGAGTCCTAGTGAG
AAGCCAGCATGGGCCAAGGAGATCCCACATCTGGTAGCTGTAAATCCCAGGGACCTGCCTTGACCGCCTCCCAG
TCAGTGACGAGCAGCCACAAGGGCCTCTCTGGGTTTCAGGAGGCTCTGAACGTGACCTCCACCGCGTGGAG
ATGCCACGCCAGAATCAGATCCCACCTCGGAAAATCCTCCTCTCCCCACTCGCATTGAAAAGTTTGACCGAAGC
TCTTGGTTACGACAGGAAGAAGACATTCCACCAAAGGTGCCTCAAAGAACAACCTTCTATATCCCAGCATTAGCC
AGAAAGAATTCTCCTGGGAATGGTAGTGCTCTGGGACCCAGACTAGGATCTCAACCCATCAGAGCAAGCAACCCCT
GATCTCCGGAGAATGAGCCCATCTTGAGAGAGCCCCCTTGAGAGGACCAGCAGTGGCAGTTTCTCCAGCTCCAGC
ACCCCTAGCTCCCAGCCCAGCTCCCAAGGAGGCTCCCAGCCTGGATCACAAAGCAGGATCCAGTGAACGCACCAGA
GTTCGAGCCAACAGTAAGTCAGAAGGATCACCTGTGCTTCCCCATGAGCCTGCCAAGGTGAAACCAGAAGAATCC
AGGGACATTACCCGGCCAGTCGACCAGCTGATCTGACGGCATTAGCCAAAGAACTAAGAGAATCCGGATTGAA
GAAACAAACCGCCCAATGAAGAAGGTGACTGATTACTCTCTCCTCAGTGAGGAGTCAGAAAGTAGCGAGGAAGAG
GAGGAAGATGGAGAGAGCGAGACCCATGATGGGACAGTGGCTGTGAGCGACATACCCAGACTGATACCAACAGGA
GCTCCAGGCAGTAACGAGCAGTACAATGTGGGATGTTGGGAGCGCATGGGCTGGAGACCTCTCATCCGGAGCAGT
TTCAGCGGCAGTATTTCAAGAGAAGGAACCTTGATGATTAGAGAGACGTCTGGAGAGAAGAAGCGATCTGGCCAC
AGTGACAGCAATGGCTTTGTGTCACATCAACCTCCCTGACCTGGTGCAGCAGAGCCATTCTCCAGCTGGAACC
CCGACTGAGGGACTGGGGCGCGTCTCAACCCATTCCCAGGAGATGGACTCTGGGACTGAATATGGCATGGGGAGC
AGCACCAAAGCCTCCTTACCCCCCTTTGTGGACCCAGAGTATACCAGACGTCTCCCACTGATGAAGATGAAGAG
GATGAGGAATCATCAGCCGAGCTCTGTTTACTAGCGAAGTTCTTAGGCAAGAACAGGCCAAACTCAATGAAGCA
AGAAAGATTTTCGGTGGTAAATGTAAACCAACCAACATTTCGGCCTCATAGCGACACACCAGAAATCAGAAAATAC
AAGAAACGATTCAACTCAGAAATACTTTGTGCAGCTCTGTGGGGTGTAACCTTCTGGTGGGGACTGAAAATGGC
CTGATGCTTTTGGACCGAAGTGGGCAAGGCAAAGTCTATAATCTGATCAACCGGAGGCGATTTAGCAGATGGAT
GTGCTAGAGGGACTGAATGTCCTTGTGACAATTTAGGAAAGAAGAATAAGCTACGAGTTTACTATCTTTCATGG
TTAAGAAACAGAATACTACATAATGACCCAGAAGTAGAAAAGAAACAAGGCTGGATCACTGTTGGGGACTTGGAA
GGCTGTATACATTATAAAGTTGTTAAATATGAAAGGATCAAATTTTGGTGATTGCCTTAAAGAATGCTGTGGAA
ATATATGCTTGGGCTCCTAAACCGTATCATAAATTCATGGCATTTAAGTCTTTTGCAGATCTCCAGCACAAAGCCT
CTGCTAGTTGATCTCACGGTAGAAGAAGGTCAAAGATTAAAGGTTATTTTGGTTTACACACTGGTTTCCATGTA
ATTGATGTTGATTCAGGAAACTCTTATGATATCTACATACCATCTCATATTACGGGCAATATCACTCCTCATGCT
ATTGTCATCTTGCTTAAACAGATGGAATGGAATGCTTGTGTTGCTATGAGGATGAGGGGGTGTATGTAAACACC
TATGGCCGGATAACTAAGGATGTGGTGCTCCAATGGGGAGAAATGCCACGTCTGTGGCTACATTCAATCCAAT
CAGATAATGGGCTGGGGCGAGAAAGCTATTGAGATCCGGTCAGTGGAACAGGACATTTGGATGGAGTATTTATG
CATAAGCGAGCTCAAAGGTTAAAGTTTCTATGTGAAAGAAATGATAAGGTATTTTTCATCCGTGCGATCTGGA
GGAAGTAGCCAAGTGTTTTTCATGACCCTCAACAGAAATTCCATGATGAACTGGTAA

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Figure 29

1 MASDSPARSLDEIDLALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIATYYGAFIKKNPPGMDDQLWLVMFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIHRDIKGQNVLLTENAEVKLVDFGVSAQLDR
181 TVGRRNTFIGTPYWMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKKWSKKFQSFIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVRI
301 QLKDHIDRTKKKRGEKDETEYEYSGSEEEEEENDSGEPSSIILNLPGESTLRRDFLRLQLA
361 NKERSEALRRQOLEQQQRENEEHKQLLAERQKRIEEQKEQRRRLLEEQQRREKELRKQOE
421 REQRRHYEEQMRREEERRRAEHEQEYKRKQLEEQRQAERLQRQLKQERDYLVSLOHQHQE
481 QRPVEKKPLYHYKEGMSPEKPAWAKEVEERSRLNRQSSPAMPHKVANRISDPNLPPRSE
541 SFSISGVQPARTPPMLRPVDPQIPHLVAVKSQGPALTASQSVHEQPTKGLSGFQEALNVT
601 SHRVEMPRQNSDPTSENPLPLTRIEKFDRSSWLROEEDIIPKVPQRTTISPALARKNSP
661 GNGSALGPRLGSQPIRASNPDLRRTPILESPLOQTSSGSSSSSSSTPSSQPSQGGSQPG
721 SQAGSSERTRVRANSKSEGSPLPHEPAKVKPEESRDITRPSRPASYKKAIDEDLTALAK
781 ELRELRIEETNRPMKKVTDYSSSSEESSESEEEEEEDGESETHDGTAVSDIPLRIPTGAP
841 GSNEQYNVGMVGTHGLETSHADSFGSISREGTLMIRETSGEKKRSGHSDSNGFAGHINL
901 PDLVQQSHSPAGTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASFTPFVDPRVYQTSPTDE
961 DEEDEESSAAALFTSELLRQEQAKLNEARKISVVNVNPTNIRPHSDTPEIRKYKKRFNSE
1021 ILCALWGVNLLVGTENGLMLLDRSGQGVYNLINRRRFQQMDVLEGLNVLVTISGKKNK
1081 LRVYYSWLRNRILHNDPEVEKKQGWITVGDLEGCIHYKVVKYERIKFLVIALKNAVEIY
1141 AWAPKPYHKFMAFKSFADLQHKPLLVDLTVEEGQRLKVI FGSHTGFHVIDVDSGNSYDIY
1201 IPSHIQGNITPHAIVILPKTDGMEMLVCYEDEGVYVNTYGRITKDVLQWGEMPTSVAYI
1261 HSNQIMGWGEKAIEIRSVETGHLDGVFMHKRAQRLKFLCERNKVFASVRSGGSSQVFF
1321 MTLNRNSMMNWZ

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Figure 30

1 MASDSPARSLDEIDL SALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIATYYGAFIKKNPPGMDDQLWLVMFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIHRDIKGQNVLLTENAEVKLVDFGVSAQLDR
181 TVGRRNTFIGTPYWMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKKWSKKFQSFIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVRI
301 QLKDHIDRTKKKRGEKDETEYEYSGSEEEEEENDSGEPSSIILNLPGESTLRRDFLRQLA
361 NKERSEALRRQQLQEQQRENEEHKRQLLAERQKRIEEQKEQRRRLEEQQRREKELRKQQE
421 REQRRHYEEQMRREEERRRAEHEQEYIRROLEEEQROLEILQQQLLHEQALLLEYKRKQL
481 EEQRQAERLQRQLKQERDYLVSLOHQROEQRPVEKKPLYHYKEGMSPEKPAWAKEI PHL
541 VAVKSQGPALTASQSVHEQPTKGLSGFQEALNVTSHRVEMPRQNSDPTSENPLPTRIEK
601 FDRSSWLQREEDIPPKVPQRTTISPALARKNSPGNGSALGPRLGSQPIRASNPDLRRT
661 PILESPLQRTSSGSSSSSSSTPSSQPSQGSQPGSQAGSSERTRVRANSKSEGSPVLPHE
721 PAKVKPEESRDITRPSRPASYKKAIDEDLTALAKELRELRIEETNRPMKKVTDYSSSSEE
781 SESSEEEEDGESETHDGTVAVSDIPRLIPTGAPGSNEQYNVGMVGTHGLETSHADSFSG
841 SISREGTLMIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSHSPAGTPTEGLGRVSTHSQ
901 EMDSGTEYGMGSSTKASFTPFDPRVYQTSPTDEDEEDEESSAAALFTSELLRQEQAKLN
961 EARKISVVNVNPTNIRPHSDTPEIRKYKKRFNSEILCAALWGVNLLVGTENGLMLLDRSG
1021 QGKVYNLINRRRFQQMDVLEGLNVLVTISGKKNKL RVYYLSWLRNRILHNDPEVEKKQGW
1081 ITVGDLGCIHYKVVKYERIKFLVIALKNAVEIYAWAPKPYHKFMAFKSFADLQHKPLL
1141 DLTVEEGQRLKVI FGSHTG FHVLDVDSGNSYDIYIPSHIQGNITPHAIVILPKTDGMEML
1201 VCYEDEGVYVNTYGRITKDVLQWGEMPTSVAYIHSNQIMGWGEKAIEIRSVETGHLDGV
1261 FMHKRAQRLKFLCERNKVFVASFVRSGGSSQVFFMTLNRNSMMNWZ

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Figure 31

1 MASDSPARSLDEIDLSALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHRNIAITYYGAFIKKNPPGMDDQLWLVMFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIHRDIKGQNVLLTENA EVKLVD FGVSAQLDR
181 TVGRRNTFIGTPYWMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKKWSKKFQSFIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVRI
301 QLKDHIDRTKKKRGEKDETEYEYSGSEEEEEENDSGEPSSIILNLPGESTLRRDFLRLQLA
361 NKERSEALRRQOLEQQORENEEHKRQLLAERQKRIEEQKEQRRRLLEEQQRREKELRKQQE
421 REQRRHYEEQMRREEERRRAEHEQEYIRROLEEEQRQLEILOQQLLHEQALLLEYKRKQL
481 EEQRQAERLQRQLKQERDYLVSLOHQHQEQRPVEKKPLYHYKEGMSPEKPAWAKEVEER
541 SRLNRQSSPAMPHKVANRISDPNLPPRSEFSISGVQPARTPPMLRPVDPQIPHLVAVKS
601 QGPALTASQSVHEQPTKGLSGFQEQALNVTSHRVEMPRQNSDPTSENPLPTRIEKFDRSS
661 WLRQEEDIPPKVPQRTTSSISPALARKNSPGNGSALGPRLGSQPIRASNPDLRRTEPILES
721 PLQRTSSGSSSSSTPSSQPSQGGSQPGSQAGSSERTRVRANSKSEGPSVLPHEPAKVK
781 PEESRDITRPSRPADLTALAKELRELRIEETNRPMMKVTDYSSSSESESESEEEEEEDGES
841 ETHDGTAVSDIPRLIPTGAPGSNEQYNVGMVGTHGLETSHADSFSGSISREGTLMIRET
901 SGEKKRSGHSDSNGFAGHINLPDLVQQSHSPAGTPTEGLGRVSTHSQEMDSGTEYGMGSS
961 TKASFTPFDPRVYQTSPTDEDEEDEESSAAALFTSELLRQEQAKLNEARKISVVNVNPT
1021 NIRPHSDTPEIRKYKKRFNSEILCAALWGVNLLVGTENGLMLLDRSGQGKVVNLINRRRF
1081 QQMDVLEGLNVLVTISGKKNKLRVYYLSWLRNRILHNDPEVEKKQGWITVGDLEGCIHYK
1141 VVKYERIKFLVIALKNAVEIYAWAPKPYHKFMAFKSFADLQHKPLLVDLTVEEGQRLKVI
1201 FGSHTGFHVIDVDSGNSYDIYIPSHIQGNITPHAIVILPKTDGMEMLVCYEDEGVYVNTY
1261 GRITKDVVLQWGEMPTSVAYIHSNQIMGWGEKAIEIRSVETGHLDGVFMHKRAQRLKFLC
1321 ERNDKVFFASVRSGGSSQVFFMTLNRNSMMNWZ

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Figure 33

1 MASDSPARSLDEIDLSALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIATYYGAFIKKNPPGMDDQLWLVMEFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIHRDIKGQNVLLTENA EVKLVD FGVSAQLDR
181 TVGRNRTFIGTPYWMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKKWSKKFQSFIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVRI
301 QLKDHIDRTKKKRGEKDETEYEYSGSEEEEEENDSGEPSSIILNLPGESTLRRDFLRLQLA
361 NKERSEALRRQQLQQQRENEEHKROLLAERQKRIEEQKEQRRRLEEQQRREKELRKQQE
421 REQRRHYEEQMRREEERRRAEHEQEYKRKQLEEQRQAERLQRQLKQERDYLVS LQHQRQE
481 QRPVEKKPLYHYKEGMS PSEKPAWAKEVEERSRLNRQSSPAMPHKVANRISDPNLPPRSE
541 SFSISGVQPARTPPMLRPVDPQIPHLVAVKSQGPALTASQSVHEQPTKGLSGFQEALNVT
601 SHRVEMPRQNSDPTSENPLPTRIEKFDRSSWL RQEEDI PP KVPQRTT SIS PALARKNSP
661 GNGSALGPRLGSQPIRASNPDLRRTEPILESPLQRTSSGSSSSSSSTPSSQPSQGGSQPG
721 SQAGSSERTRVRANSKSEGSPVLPHEPAKVKPEESRDITRPSRPADLTALAKELRELRIE
781 ETNRPMKKVTDYSSSSEESSESEEEEEEDGESETHDGTAVASDIPRLIPTGAPGSNEQYNV
841 GMVGTHGLETSHADS FSGSISREGTLMIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSH
901 SPAGTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASFTPFDPRVYQTSPTDEDEEDEESS
961 AAALFTSELLRQEQA KLNEARKISVVNVNPTNIRPHSDTPEIRKYKKRFNSEILCAALWG
1021 VNLLVGTENGLMLLDRSGQGKVYNLINRRRFQQMDVLEGLNVLVTISGKKNKLRVYYLSW
1081 LRNRILHNDPEVEKKQGWI TVGDLEGC IHYKVVKYERIKFLVIALKNAVEIYAWAPKPYH
1141 KFMAFKSFADLQHKPLLVDLTVEEGQRLKVI FGSHTGFHVIDVD SGNSYDIYIPSHIQGN
1201 ITPHAIVILPKTDGMEMLVCYEDEGVYVNTYGRITKD VVLQWGEMPTSVAYIHSNQIMGW
1261 GEKAI EIRSVETGHL DGVFMHKRAQRLKFLCERN DKVFFASVRSGGSSQVFFMTLNRNSM
1321 MNWZ

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Figure 34

1 MASDSPARSLDEIDLSALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIATYYGAFIKKNPPGMDDQLWLVMFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIHRDIKGQNVLLTENA EVKLVD FGVSAQLDR
181 TVGRRNTFIGTPYWMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKKWSKKFQSFIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVRI
301 QLKDHIDRTKKKRGEKDETEYEYSGSEEEEEENDSGEPSSI LNLPGESTLRRDFLRLQLA
361 NKERSEALRRQQL EQQRENEEHKRQLLAERQKRIEEQKEQRRRL EEQQRREKELRKQQE
421 REQRRHYEQMRREEERRRAEHEQEYIRRQLEEEQRQLEILQQQLLHEQALLLEYKRRQL
481 EEQRQAERLQRQLKQERDYLVS LQHQRQEQR PVEKKPLYHYKEGMS PSEKPAWAKEI PHL
541 VAVKSQGPALTASQSVHEQPTKGLSGFQEALNVTSHRVEMPRQNSDPTSENPPLPTRIEK
601 FDRSSWL RQEEDI PPKV PQR TTSISPALARKNSPGNGSALGPRLGSQPIRASNPDLRRT E
661 PILESPLQRTSSGSSSSSSTPSSQPSQGGSQPGSQAGSSERTRVRANSKSEGSPVLPHE
721 PAKVKPEESRDITRPSRPADLTALAKELRELRIEETNRPMKKVTDYSSSSEESSESESESE
781 EDGESETHDGTVAVSDI PRLIPTGAPGSNEQYNVGMVGTHGLETSHADSFSGSISREGTL
841 MIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSHSPAGTPTEGLGRVSTHSQEMDSGTEY
901 GMGSSTKASFTPFDPRVYQTSPTDEDEDEEESAAALFTSELLRQEQA KLNEARKISV
961 NVNPTNIRPHSDTPEIRKYKRFNSEILCAALWGVNLLVGTENGLMLLDRSGQGKVYNLI
1021 NRRRFQQMDVLEGLNVLVTISGKKNKLRVYYLSWLRNRILHNDPEVEKKQGWITVGDLEG
1081 CIHYKVVKYERIKFLVIALKNAVEIYAWAPKPYHKFMAFKSFADLQHKPLLVDLTVEEGQ
1141 RLKVIFGSHTGFHVIDVDSGNSYDIYIPSHIQGNITPHAIVILPKTDGMEMLVCYEDEGV
1201 YVNTYGRITKDVVLQWGEMPTSVAYIHSNQIMGWGEKAIEIRSVETGHLDGVMHKRAQR
1261 LKFLCERNDKVFFASVRSGSSQVFFMTLNRNSMMNWZ

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